

QUENCE LISTING

<110> Huse, William D. Freedman, Michael H.



<120> Method for Identifying Optimal Binding Ligands to a Receptor

<130> P-IX 3280

<140> US 09/169,048

<141> 1998-10-08

<150> 60/112,011

<151> 1997-10-09

<160> 28

<170> PatentIn Ver. 2.1

<210> 1

<211> 24

<212> DNA

<213> Mus musculus

<220>

<221> CDS

<222> (1)..(24)

<400> 1

agc tca agt gta agt ttc atg aac Ser Ser Ser Val Ser Phe Met Asn 1 5

<210> 2

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<212> PRT

<213> Mus musculus

<400> 2

Ser Ser Ser Val Ser Phe Met Asn 1 5

<210> 3

<211> 24

24

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                                                                    24
Ser Ser Ser Val Arg Phe Met Asn
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<210> 5
<211> 24
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Ser Glu Ser Val Asn Leu Met Asn
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Ser Ser Ser Val Asn Phe Met Asn
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<211> 24

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Ser Ser Thr Val Ser Phe Met Asn
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Ser Ser Ser Val Ala Tyr Met Asn
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Ser Gln Ser Ala Lys His Met Asn
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Ala Thr Ser Asn Leu Ala Ser Gly
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Ala Thr Ser Asn Leu Ala Ser Gly
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                 5
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Ala Thr Glu Lys Leu Ala Ser Gly
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construct

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<400> 20
Ala Thr Val Asn Leu Ala Ser Gly
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<222> (1)..(24)

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Ala Thr Val Asn Leu Ala Ser Gly
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Ala Thr Val Asn Leu Ala Ser Gly
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<210> 23
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gcc aca tcc agg gcg gct tct gga
Ala Thr Ser Arg Ala Ala Ser Gly
 1
                 5
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<213> Artificial Sequence
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<223> Description of Artificial Sequence: synthetic

construct

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<220> <221> CDS

<222> (1)..(24)

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Ala Thr Ser Arg Ala Ala Ser Gly
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Ala Thr Gln Asn Leu Ala Ser Gly
1
                  5
<210> 26
<211> 8
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24

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<220>
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<400> 27
gcc aca tcc aat ttg gct tct gga
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Ala Thr Ser Asn Leu Ala Ser Gly
                  5
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Ala Thr Ser Asn Leu Ala Ser Gly
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